## SEQUENCE LISTING

<110> Elliott, Steven G. Rogers, Norma Busse, Leigh Anne <120> G-Protein Coupled Receptor Molecules and Uses Thereof <130> 02-076 <140> <141> <150> 60/269,040 <151> 2001-02-14 <160> 22 <170> PatentIn Ver. 2.0 <210> 1 <211> 1038 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(1038) atg tac aac ggg tcg tgc tgc cgc atc gag ggg gac acc atc tcc cag Met Tyr Asn Gly Ser Cys Cys Arg Ile Glu Gly Asp Thr Ile Ser Gln gtg atg ccg ccg ctc att gtg gcc ttt gtg ctg ggc gca cta ggc Val Met Pro Pro Leu Leu Ile Val Ala Phe Val Leu Gly Ala Leu Gly 20 aat ggg gtc gcc ctg tgt ggt ttc tgc ttc cac atg aag acc tgg aag Asn Gly Val Ala Leu Cys Gly Phe Cys Phe His Met Lys Thr Trp Lys ccc agc act gtt tac ctt ttc aat ttg gcc gtg gct gat ttc ctc ctt 192 Pro Ser Thr Val Tyr Leu Phe Asn Leu Ala Val Ala Asp Phe Leu Leu 55 50 atg atc tgc ctg cct ttt cgg aca gac tat tac ctc aga cgt aga cac 240 Met Ile Cys Leu Pro Phe Arg Thr Asp Tyr Tyr Leu Arg Arg Arg His 70 65 288 tgg gct ttt ggg gac att ccc tgc cga gtg ggg ctc ttc acg ttg gcc Trp Ala Phe Gly Asp Ile Pro Cys Arg Val Gly Leu Phe Thr Leu Ala 85 atg aac agg gcc ggg agc atc gtg ttc ctt acg gtg gtg gct gcg gac 336 Met Asn Arg Ala Gly Ser Ile Val Phe Leu Thr Val Val Ala Ala Asp 105

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His	Gly	Ala	Leu 260		Ile	Thr	Leu	Ser 265	Phe	Thr	Tyr	Met	Asn 270	Ser	Met	816
Leu	Asp	275	Leu	gtg Val	Tyr	Tyr	Phe 280	Ser	Ser	Pro	Ser	Phe 285	Pro	Lys	Phe	864
Tyr	290	Lys )	Leu	Lys	Ile	295	Ser	Leu	Lys	Pro	300	Gln	Pro	Gly		912
Ser 305	Lys	Thr	Glr	Arg	310	Glu	Glu	Met	Pro	315	e Ser	Asn	. Leu	. Gly	cgc Arg 320	960
agg Arg	g agt g Ser	tgo Cys	ato Ile	agt Ser 325	Val	gca . Ala	aat Asr	agt Ser	Phe 330	e Glr	ago Ser	cag Glr	f tct Ser	gat Asp 335	Gly 9gg	1008

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<213> Homo sapiens

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Pro Ser Thr Val Tyr Leu Phe Asn Leu Ala Val Ala Asp Phe Leu Leu 50 55 60

Met Ile Cys Leu Pro Phe Arg Thr Asp Tyr Tyr Leu Arg Arg His
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Trp Ala Phe Gly Asp Ile Pro Cys Arg Val Gly Leu Phe Thr Leu Ala 85 90 95

Met Asn Arg Ala Gly Ser Ile Val Phe Leu Thr Val Val Ala Ala Asp 100 105 110

Arg Tyr Phe Lys Val Val His Pro His His Ala Val Asn Thr Ile Ser 115 120 125

Thr Arg Val Ala Ala Gly Ile Val Cys Thr Leu Trp Ala Leu Val Ile 130 135 140

Leu Gly Thr Val Tyr Leu Leu Leu Glu Asn His Leu Cys Val Gln Glu 145 150 155 160

Thr Ala Val Ser Cys Glu Ser Phe Ile Met Glu Ser Ala Asn Gly Trp 165 170 175

His Asp Ile Met Phe Gln Leu Glu Phe Phe Met Pro Leu Gly Ile Ile 180 185 190

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195 200 205

Leu Ala Arg Gln Ala Arg Met Lys Lys Ala Thr Arg Phe Ile Met Val 210 215 220

Val Ala Ile Val Phe Ile Thr Cys Tyr Leu Pro Ser Val Ser Ala Arg 225 230 235 240

Leu Tyr Phe Leu Trp Thr Val Pro Ser Ser Ala Cys Asp Pro Ser Val

245 250 255

His Gly Ala Leu His Ile Thr Leu Ser Phe Thr Tyr Met Asn Ser Met 260 265 270

Leu Asp Pro Leu Val Tyr Tyr Phe Ser Ser Pro Ser Phe Pro Lys Phe 275 280 285

Tyr Asn Lys Leu Lys Ile Cys Ser Leu Lys Pro Lys Gln Pro Gly His 290 295 300

Ser Lys Thr Gln Arg Pro Glu Glu Met Pro Ile Ser Asn Leu Gly Arg 305 310 315 320

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Met Pro Val

ctc tct cca act gct atg gac aac ggg tcg tgc tgt ctc atc gag ggg 406 Leu Ser Pro Thr Ala Met Asp Asn Gly Ser Cys Cys Leu Ile Glu Gly

gag ccc atc tcc cag gtg atg cct cct cta ctc atc ctg gtc ttc gtg 454
Glu Pro Ile Ser Gln Val Met Pro Pro Leu Leu Ile Leu Val Phe Val
20 25 30 35

ctt ggc gcc ctg ggc aac ggc ata gcc ctg tgc ggc ttc tgc ttt cac 502 Leu Gly Ala Leu Gly Asn Gly Ile Ala Leu Cys Gly Phe Cys Phe His 40 45 50

atg Met	aag Lys	acc Thr	tgg Trp 55	aag Lys	tca Ser	agc Ser	act Thr	att Ile 60	tac Tyr	ctt Leu	ttc Phe	aac Asn	ttg Leu 65	gct Ala	gtg Val	550
											cgg Arg					598
ctc Leu	aga Arg 85	cgc Arg	aga Arg	caĉ His	tgg Trp	att Ile 90	ttt Phe	gga Gly	gat Asp	atc Ile	gcc Ala 95	tgt Cys	cgc Arg	ctg Leu	gtc Val	646
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tcg aac ctc tgc agt aag agc tcc atc gat ggg gca aat cgt tcc cag 1: Ser Asn Leu Cys Ser Lys Ser Ser Ile Asp Gly Ala Asn Arg Ser Gln 325 330 335	366
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35 40 45

Cys Phe His Met Lys Thr Trp Lys Ser Ser Thr Ile Tyr Leu Phe Asn 50 55 60

Leu Ala Val Ala Asp Phe Leu Leu Met Ile Cys Leu Pro Leu Arg Thr 65 70 75 80

Asp Tyr Tyr Leu Arg Arg Arg His Trp Ile Phe Gly Asp Ile Ala Cys
85 90 95

Arg Leu Val Leu Phe Lys Leu Ala Met Asn Arg Ala Gly Ser Ile Val

Phe Leu Thr Val Val Ala Val Asp Arg Tyr Phe Lys Val Val His Pro 115 120 125

His His Met Val Asn Ala Ile Ser Asn Arg Thr Ala Ala Ala Thr Ala 130 135 140 Cys Val Leu Trp Thr Leu Val Ile Leu Gly Thr Val Tyr Leu Leu Met 160 145 Glu Ser His Leu Cys Val Gln Gly Thr Leu Ser Ser Cys Glu Ser Phe 165 170 Ile Met Glu Ser Ala Asn Gly Trp His Asp Val Met Phe Gln Leu Glu 185 Phe Phe Leu Pro Leu Thr Ile Ile Leu Phe Cys Ser Val Asn Val Val 195 Trp Ser Leu Arg Arg Arg Gln Gln Leu Thr Arg Gln Ala Arg Met Arg 215 Arg Ala Thr Arg Phe Ile Met Val Val Ala Ser Val Phe Ile Thr Cys Tyr Leu Pro Ser Val Leu Ala Arg Leu Tyr Phe Leu Trp Thr Val Pro 245 250 Thr Ser Ala Cys Asp Pro Ser Val His Thr Ala Leu His Val Thr Leu Ser Phe Thr Tyr Leu Asn Ser Met Leu Asp Pro Leu Val Tyr Tyr Phe 275 280 Ser Ser Pro Ser Leu Pro Lys Phe Tyr Ala Lys Leu Thr Ile Cys Ser 295 Leu Lys Pro Lys Arg Pro Gly Arg Thr Lys Thr Arg Arg Ser Glu Glu 315 310 Met Pro Ile Ser Asn Leu Cys Ser Lys Ser Ser Ile Asp Gly Ala Asn 325

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				_	aag Lys			_	_	_						553
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					aga Arg 85											649
					ttc Phe											697
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- Cys Phe His Met Lys Thr Trp Lys Ser Ser Thr Ile Tyr Leu Phe Asn 50 55 60
- Leu Ala Val Ala Asp Phe Leu Leu Met Ile Cys Leu Pro Leu Arg Thr 65 70 75 80
- Asp Tyr Tyr Leu Arg Arg Arg His Trp Ile Leu Gly Asp Ile Pro Cys
  85 90 95
- Arg Leu Val Leu Phe Met Leu Ala Met Asn Arg Ala Gly Ser Ile Val 100 105 110
- Phe Leu Thr Val Val Ala Val Asp Arg Tyr Phe Lys Val Val His Pro 115 120 125
- His His Met Val Asn Ala Ile Ser Asn Arg Thr Ala Ala Ile Val 130 135 140
- Cys Val Leu Trp Thr Leu Val Ile Leu Gly Thr Val Tyr Leu Leu Met 145 150 155 160
- Glu Ser His Leu Cys Val Arg Gly Met Val Ser Ser Cys Glu Ser Phe 165 170 175
- Ile Met Glu Ser Ala Asn Gly Trp His Asp Ile Met Phe Gln Leu Glu 180 185 190
- Phe Phe Leu Pro Leu Thr Ile Ile Leu Phe Cys Ser Phe Lys Val Val 195 200 205
- Trp Ser Leu Arg Gln Arg Gln Gln Leu Thr Arg Gln Ala Arg Met Arg 210 215 220
- Arg Ala Thr Arg Phe Ile Met Val Val Ala Ser Val Phe Ile Thr Cys 225 230 235 240
- Tyr Leu Pro Ser Val Leu Ala Arg Leu Tyr Phe Leu Trp Thr Val Pro 245 250 255
- Ser Ser Ala Cys Asp Pro Ser Val His Ile Ala Leu His Val Thr Leu 260 265 270
- Ser Leu Thr Tyr Leu Asn Ser Met Leu Asp Pro Leu Val Tyr Tyr Phe 275 280 285
- Ser Ser Pro Ser Phe Pro Lys Phe Tyr Ala Lys Leu Lys Ile Arg Ser 290 295 300
- Leu Lys Pro Arg Arg Pro Gly Arg Ser Gln Ala Arg Arg Ser Glu Glu 305 310 315 320

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